GenCore version 4.5

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OM nucleic - nucleic search, using sw model

Run on: November 8, 2001, 12:59:21 ; Search time 101.15 Seconds
(without alignments)
13812.583 Million cell updates/sec

Title: US-09-227-881-3
Perfect score: 6169
Sequence: 1 atctttgttcagttacctc.........cttgtgccctccatgtcag 6169

Scoring table: OLIGO_NUC
Gapop 60.0 , Gapext 60.0

Searched: 351203 seqs, 113238999 residues

Word size : 8

Total number of hits satisfying chosen parameters:　540559

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : Issued_Patents_NA:*
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB ID	Description
1	6169	100.0	6169	4	US-08-938-669A-3	Sequence 3, Appl1
2	5008	81.2	5300	4	US-08-938-669A-1	Sequence 1, Appl1
3	4476	72.6	5304	4	US-08-938-669A-2	Sequence 2, Appl1
4	640	10.4	1548	4	US-08-938-669A-26	Sequence 26, Appl1
5	640	10.4	1999	2	US-08-645-900A-2	Sequence 2, Appl1
6	640	10.4	1999	2	US-08-882-238A-2	Sequence 2, Appl1
7	640	10.4	1999	2	US-08-667-790A-2	Sequence 2, Appl1
8	640	10.4	1999	3	US-09-220-459-2	Sequence 2, Appl1
9	640	10.4	1999	2	US-09-220-459-2	Sequence 2, Appl1
10	604	9.8	1512	2	US-08-546-568B-2	Sequence 2, Appl1
11	604	9.8	1512	2	US-08-645-900A-3	Sequence 3, Appl1
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16	553	9.0	1512	4	US-08-822-999-2	Sequence 2, Appl1
17	538	8.7	1985	2	US-08-791-347-1	Sequence 1, Appl1
18	527	8.5	1509	2	US-08-791-347-2	Sequence 2, Appl1
19	366	5.9	1968	1	PCT-US95-14024-2	Sequence 2, Appl1
20	366	5.9	1969	1	US-08-336-235A-2	Sequence 2, Appl1
21	366	5.9	1969	1	US-08-336-235A-2	Sequence 2, Appl1
22	330	5.3	1491	1	US-08-649-432-3	Sequence 3, Appl1
23	330	5.3	1491	1	US-08-336-235A-3	Sequence 3, Appl1
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26	88	1.4	643	2	US-08-822-999-9	Sequence 9, Appl1
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41	36	0.6	2713	6	US-08-916-901-6	Sequence 6, Appl1
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43	36	0.6	36741	4	US-09-301-665-3	Sequence 3, Appl1
44	36	0.6	246240	2	US-08-724-394A-20	Sequence 20, Appl1
45	36	0.6	246240	2	US-08-724-394A-21	Sequence 21, Appl1

ALIGNMENTS

RESULT 1
US-08-938-669A-3
; Sequence 3, Application US/0893866A
; Patent No. 6171788
; GENERAL INFORMATION:
; APPLICANT: Nguyen, Thai D.
; APPLICANT: Polansky, Jon R.
; TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS,
; TITLE OF INVENTION: PROGNOSIS AND TREATMENT OF GLAUCOMA AND
; TITLE OF INVENTION: RELATED DISEASES
; NUMBER OF SEQUENCES: 32
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Howrey & Simon
; STREET: 1299 Pennsylvania Avenue, N.W.
; CITY: Washington
; STATE: DC
; COUNTRY: USA
; ZIP: 20004-2402
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSEQ for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/938,669A
; FILING DATE:
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/791,154
; FILING DATE: 28-JAN-1997
; ATTORNEY/AGENT INFORMATION:
; NAME: Mendelson, Elliot
; REGISTRATION NUMBER: P-42,878
; REFERENCE/DOCKET NUMBER: 07425-0034
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202 383-6857
; TELEFAX: 202 383-6610
; TELEX:
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 6169 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-938-669A-3

Query Match	100.0%;	Score 6169;	DB 4;	Length 6169;
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Matches 6169;	Conservative	0;	Mismatches	0;	Indels	0;	Gaps	0;

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Qy  3301 ggtagctgtttgcctggcaattcaaaaaactggtgccagagagagcaagtggaaaatgccagagattg 3360
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  3301 GGTAGCTTTTTTGCCCTGGCCATTCAAAAAACTGGGCCCAGAGCAGCAAGTGGAAAATGCCAGAGATTG 3360

Qy  3361 ttaaacttttcacccctgaccagcaccccaccccacggccagctcagcagctgacagccacgg 3420
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  3361 TTAAACTTTTCACCCTGACCACCACCCCACCCCACGGCCAGCTCAGCCAGCTGACAGCCACGG 3420

Qy  3421 agtgacctgcagcgcagggggaggggaggagagagagatagtgtatgaggaag 3480
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  3421 AGTGACCTGCAGCGCGCAGGGGGAGGGGAGGAGAGGGATAGTTAGTGTATGAGGCAAGAAAG 3480

Qy  3481 acagattcaatccatccaaaggggcagtgggaattgaccacacaggatatatagccagtaccctg 3540
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  3481 ACAGATTCATTCAAGGGGCAGTGGGGAATTGACCACACAGGGATATAGTCCACGTGATCCTGG 3540

Qy  3541 gttctaggaggcagggctatattgtggggggaaaaatcagtcacgacaaggagtcgggga 3600
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  3541 GTTCTAGGAGGCCAGGGCTATATTGTGGGGAAAAATCAGTTCAAGGGAAGTCGGGGA 3600

Qy  3601 cctgattctaatactatatatttttccttacaagctggtaatcctctgacacaag 3660
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  3601 CCTGATTTCTAATACTATATATTTTTCCTTTACAAGCTGGTAATTCTGACACAAG 3660

Qy  3661 gtagtaactgagggctgtaagattaactttttatctggagaaatggagaaacctctatgcactctg 3720
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Db  3661 GTAGTAACTGAGGGCTGTTAAGATTAACTTTTCCCTTTATGAGAAATGGAGAGAACCCTTATGCACTCTG 3720

Qy  3721 ggagttagcagcagggcaatcccgttctttttaacagaagaaaacattcctaaag 3780
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Db  3721 GGAGTTAGCAGCAGGGCAATCCCGTTTCTTTTAACGGAAGAGAAAATCCTAAGAG 3780

Qy  3781 taaagtccaaacaagaattcaagcctaggtcctgctgctagtaatgaattggttttgaaaaat 3840
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Db  3781 TAAAGCCAAAACAAGAATTCTAAATACTTATATTTTTTCCTTTACAAGCTGAGTAATTCTGACAAAT 3840

Qy  3841 catttcacgtcggatgttactactctgattcaagaaaatggaactagtaccctttcagctg 3900
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Db  3841 CATTTCCAGCGATGTTTAATCCTGATTTCAGAGAATTAGTTAGTTACCTCCTTTGGTCAGCTG 3900

Qy  3901 taaacaaacacccagtgtaaatgtctccaagtcaagcttaaactgacacagttcaaaa 3960
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Qy  3961 agaatagaaatcctttagagcaatgttgttctccccaatctggagctgagggc 4020
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Db  3961 AGAATAGAAATCTTTAGAGCAATCATTATTTTTCTCCCCAATCTGGAGGTGCCAGGGC 4020

Qy  4021 agtttggaaaatattttaacttacacttgttgtggttataacacataaagt 4080
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Db  4021 AGTTTGGAAAATATTTACTTCACAGTATTGACACTGTTGTGGTTATAACACTAAAGT 4080

Qy  4081 tgcttcaaaggcaatcattatttatttcaagtggcttaaagttacttctgacaagttttgtatat 4140
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Db  4081 TAAACAAACACCCAGTTGTAAATGTCTCCAAGTTCAGGCTTAACTGACAGTTAACTGCAAATCAAAA 4140

Qy  4141 ttattggctattgccatttgctttttgtttttttttttattaaatgtaaagcag 4200
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  4141 TTATTGGCTATTGCCATTTGCTTTTTGTTTTTTTTTTTATTAAATGTAAAGCAG 4200

Qy  4201 ggattaattaacctacagtccagagaaatttgaaatttgtttctctggttttaatgtaaacacatttt 4260
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  4201 GGATTAATTAACCTACAGTCCAGAGAAAATATTACTTCACAGTATTGACACTGTTGTGGTTATAACATTTT 4260

Qy  4261 tgttcttaccacccttctaaacattttacattttattccattttatttacattttcct 4320
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Db  4261 TGTTTTTTACCACCTTCTAAACACCTTCTAACATTTAATCGAATAGAGCCCATAAA 4320

Qy  4321 ctcaaagtggtaataaccagtaccctgaatttgtcattaccaatagaaaatacagacatt 4380
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Db  4321 CTCAAAGTGGTAATAACCAGTACCTGTGATTTTGTGTCATTACCAATAGAAAATCACAGACATT 4380
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Qy  4381 tatactactattacagttgttcagatacgttgtaagtgaaataatttatactcaaaacta 4440
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Db  4381 TTTATACTATTATTACAGTTGTTGCAGATACGTTGTAAGTCGAAATATTTATACTCAAAACTA 4440

Qy  4441 ctttgaaattagaccctcctgctgctggaatctcttgttcttaaacatattaaataaaac 4500
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Db  4441 CTTTGAAATTAGACCCTCCTGCTGCTGGAATCTGTTGTTTAACATATTAATAAAACATGTTTTAAA 4500

Qy  4501 atttgatatcattatcattatttgttttccttgttaatctatattt 4560
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Db  4501 ATTTTGATATTTTGATAATCATATTTGTTTTCCTTGTTAATCTATATTTT 4560

Qy  4561 atatattttgaaaacatcttctgtgagaagaggttcccaatgaggttcctgg 4620
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Db  4561 ATAATATTTGAAAACATCTTTCTGAGAAGAGGTTCCCAGATTTCACCAATGAGGTTCTTGG 4620

Qy  4621 catgcacacacacaagtaagaactgattttgagggctaaacattggttgcctgaga 4680
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  4621 CATGCACACACACACAGTAAGAACTGATTTTGAGGGCTAAACATTGGTTGCCTGACA 4680

Qy  4681 tgccaagactgaaattagaaagttcctcccaaagataacacagttgttttaaagctaggggtg 4740
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  4681 TGCCAAGACTGAAATTAGAAAGTTCTCCCAAAGATACACAGTTGTTTTTAAAGCTAGGGGTG 4740

Qy  4741 aggggggaaatctgccggcttctataggaatgctctccctggactgtgctctgtc 4800
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Db  4741 AGGGGGGAAATCTGCCGGCTTCTATAGGAATGCTCTCCCTGGACCTGTGCTCTGTC 4800

Qy  4801 cttgtgttctgggttgggctgttattttctctctgctacgtgtctaaaagactgttt 4860
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  4801 CTTGTGTTCTGGGTGGGCTGTTATTTTCTCTGTTCCTACCTCTTAAAGGACTTGTTT 4860

Qy  4861 ggatccagttcctagcgatagtgcctggccagttctcaatgagttcagg 4920
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Db  4861 GGATCTCCAGTTCCTAGCATAGTGCCTGGCACAGTTCTCAATGAGTTTGCAGAG 4920

Qy  4921 tgaatggaaatataacctagaaatatctccttgtctgaaatcagacacacagtagtcctg 4980
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Db  4921 TGAATGGAAATAAACTAGAATATCCTTGTTGAAATCAGACACACAGTAGTCCTGG 4980

Qy  4981 tgtaagtgtgtacgtgtgtgtgtgtgtggagata 5040
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Db  4981 TGTAAGTGTGTACGTGTGTGTGTGTGTGTGGAGATA 5040

Qy  5041 taggaactatttattggggtatgggtgcataaaatttgggatgtctttttaaaaagaaactc 5100
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Db  5041 TAGGAACTATTATTGGGGTGCATAAAAATTTGGGATGTTCTTTTTAAAAAGAAACTC 5100

Qy  5101 caaacagactctggaaggttattttctataggaatcttgctcggcagacgtgaaggcaaccc 5160
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Db  5101 CAAACAGACTTCTGGAAGGTTATTTTCTAAGAATCTTGCTCGGCAGACGTGAAGGCAACCC 5160

Qy  5161 cctgtgcacagcccacccaccctctcacgtggccacctctgtctcccccacatgaagggctg 5220
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Db  5161 CCTGTGCACAGCCCACCCACCCTCACGTGGCCACCTCTGTCTCCCCATGAAGGGCTG 5220

Qy  5221 gctcccagtatataaacctctctggagctcgggcggcaagggccaccaatc 5280
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Db  5221 CAGAGGAGGCCCAGCCCCATGTCAGCTCAATCCATAAACTTACAGAGATGGCAGGCAT 5280

Qy  5281 caggcaccctcagcacagagaggagagagcctcaccaagccctctgcaatga 5340
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Db  5281 CAGGCACCCTCAGCACAGCAGAGCTTTCCAGGAGGAAGCCTCACCAAGCCCTGCAATGA 5340

Qy  5341 ggtcctctgcaagttgtcagctttgggccctgaggatgtgtcagctttgggcctttgggcctttgggcct 5400
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Qy  5401 ttctggcctcgcctggttgtgggatgtgggggccaggacagctcaagctcaaggccaatg 5460
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Db  5401 TTCTGGCCTCGCCTGGTGTGTGGGATGCGGGGGCCAGGACAGCTTCAGCTCAAGGAAGGCCAATG 5460
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Qy  5461 accagaagtggccagatgccagtataccttcagtgtggccagtcccagtgaatccagctgcc 5520
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Db  5461 ACCAGAGTGGCCGGCGGGATGCCAGCCAGTATAACCTTCAGTGTGGCCAGTCCCAGTGAATCCAGCTGCC 5520

Qy  5521 cagagcaagagagccaagccaatgtcagtcaatccaataactttaccgtgggcccaccaac 5580
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Db  5521 CAGAGCAGAGCGCCAGCCAGTCAGTCAATCCATAACTTTACAGAGATACAGCAGCACCAAC 5580

Qy  5581 gcttagaacttggaggccaccaaaagctcgactcagctcccctggaagagctcctccacca 5640
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Db  5581 GCTTAGACCTGGAGCGCCACCAAAAGTCGACTCAGTTCCCTGGAGAGCTCCTCCAAT 5640

Qy  5641 tgaccttggaccaggctgcaggtcgccagagaaggttcctgg 5700
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Db  5641 TGACCTTGGACCAGGCTGCCAGGACCCCCAGGAGACCGAGCTGCACAGGAGCTGG 5700

Qy  5701 gcaccctgaggcgggagagcggaggaaccagctgaaaaccaccagagaggaggaaaggtt 5760
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Db  5701 GCACCCTGAGGCGGGAGCGGGGACCAGCTGAAAACCACCAAACCAGGTCCCCACCAGTTGGGACTGCCT 5760

Qy  5761 acagcaaacctcctccggacacaagtcagtctctggaggaggtgatatggctcgatgagg 5820
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Db  5761 ACAGCAAACCTCCTCCGAGACAAGTCAGTTCGTCTGGAGGAGAAGAACGACTAAGGCAAG 5820

Qy  5821 aaaatgagaatctggccagagaggaggttggaaagcacagccagaggagagtggagagaa 5880
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Db  5821 AAAATGAGAATCTGGCCAGAGGAAGCAGCCAGGGGTAGCAAAGGCTTGCAAAGGCTGAGAA 5880

Qy  5881 ggggcaggtgtccccagaaccccgagacacgtctcgggtcgtcaccaaggtcccagagag 5940
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Db  5881 GGGGCCAGTGTCCCCAGACCCGAGACACTGCTCCGGGTCTCCTCCCAGCCTCAGAGAAG 5940

Qy  5941 gtaagaaatgcaagagtggggggaactctgagtccagagtgatatggtctgagtagtgtgatcg 6000
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Db  5941 GTAAGCAAATGCAGAGTGGGGGGACTTCGAGTTCCAGGGTGATATGGCTCGTGAGTGACCTG 6000

Qy  6001 ctacaaggcgctccaggccctccccctgcccctttctctcctagaggactgacagctgacagac 6060
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Db  6001 CTACAGGGCGCTCCAGGCCCTCCAGGCCTGAGTTCAGCAGGTGATATGGTCTGGTAGTCCTG 6060

Qy  6061 agatgaattaaggagaacacacgatcacctccaagtattactagtattactgagtaatttagctcctgag 6120
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Db  6061 AGATGAATTAAGGAGAAAGCACACGATCACCTTCAAGTATTACTAGTAATTTAGCTCCTGAG 6120

Qy  6121 agcttcattagattagttggttcaagagttcttgtgccctccatgtcag 6169
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Db  6121 AGCTTCATTTAGATTAGTGGTTCAGAGTTCTTGTGCCCCTCCAAGTCAG 6169
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  APPLICATION NUMBER: US/08/938,669A
  FILING DATE:
  CLASSIFICATION: 435
  PRIOR APPLICATION DATA:
  APPLICATION NUMBER: 08/791,154
  FILING DATE: 28-JAN-1997
  ATTORNEY/AGENT INFORMATION:
  NAME: Mendelson, Elliot
  REGISTRATION NUMBER: P-42,878
  REFERENCE/DOCKET NUMBER: 07425-0034
  TELECOMMUNICATION INFORMATION:
  TELEPHONE: 202 383-6857
  TELEFAX: 202 383-6610
  TELEX:
  INFORMATION FOR SEQ ID NO: 1:
  SEQUENCE CHARACTERISTICS:
  LENGTH: 5300 base pairs
  TYPE: nucleic acid
  STRANDEDNESS: single
  TOPOLOGY: linear
  US-08-938-669A-1
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Query Match          81.2%;  Score 5008;  DB 4;  Length 5300;
Best Local Similarity  99.9%;  Pred No. 0;
Matches 5298;  Conservative 0;  Mismatches 1;  Indels 2;  Gaps 2;
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Qy     1 atcttgttcagttcacctcagggctattatgtatgaaatgaaataggataaccaatgtgaaag 60
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Db     1 ATCTTGTTCAGTTTACCTCAGGGCCTATTATGTATGAAATGAAATGAGATAACCAATGTGAAAG 60

Qy    61 tcctatataaactgtatagcctccaattcggatgtatgtctttgggcaggatgataaagaatca 120
         ||||||| |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    61 TCCTATAAACTGTATAGCCTCCAATTCGGATGTATGTCTTTGGGCAGGATGATAAAGAATCA 120

Qy   121 ggaagaaggagtatccacgttagccaagtcgtgtctgctcttattcttagtga 180
         |||||||||||||||||||||||||| |||||  |||||||||||   |||||
Db   121 GGAAGAAGGAGTATCCACGTTAGCCAAGTCGTCAGGCCTCGTCTGCTCTTATTTTAGTGA 180

Qy   181 cagatgttgctcctgacagaggaagctatctctcaggaaacatcacatccaatatggtaaatc 240
         |||||||||||||||||| ||||||| || ||||||||||||||||||||||||||||||||||
Db   181 CAGATGTTGCTCCTGACAGAAGCTATTCTTCAGGAAACATCACATCCAATATGGTAAATC 240

Qy   241 catcaaacacggaggctaagaaacaggaatgagagtgggcacttgcccaaggagaaaatgccag 300
         ||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   241 CATCAAACAGGAGCTAAGAAACAGGAATGAGAGTGGGCACTTGCCCAAGGAGAAAATGCCAG 300

Qy   301 gagggcaaatatgatgaaaataaaactttcccttgttttaattccaggaaaatg 360
         || | |||| ||||||||||||||||||||||| ||||||||| |||||||| |
Db   301 GAGAGCAAATAATGATGAAAATAAAACTTTCCCTTTGTTTTAATTTCAGGAAAAATG 360

Qy   361 atgaggaccaaaatcaatgaataggaaaagacgctcagaaaaagaaagatgtttccaaattgg 420
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Db   361 ATGAGGACCAAAATCAATGAATAGGAAAAGACAGCTCAGAAAAACAGTGTTCCAAATTGG 420

Qy   421 taattaagtgtattgttccttgggaagaagagacctccatgtgagctgatgggaaaatgggaa 480
         ||||||||| || |||||||||||||| || ||||||||||||  |  ||||| |||||||||
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Qy   481 aaacgtcaaaagcatgactgatcagaatcccaaagtggattatttattcttaaaaacccagat 540
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Db   481 AAACGTCAAAAGCATGATCAGATCCCAAAGTGGATTATTTATTCTTAAAAACCAGAT 540

Qy   541 ggcatcactctggggaggcaagttcagggagtcatgttagcaaaggacataacaatac 600
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Db   541 GGCATCACTCTGGGGAGGCAAGTTCAGGGACTGGAATGGTCATGTTAGCAAAGGACATAACAATAC 600

Qy   601 agcaaaatcaaaattccgcaaatgcaggagaaaatgggaacttcataac 660
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Db   601 AGCAAAATCAAAATTCCGCAAATGCAGGAGAAAATGGGAACTTCATAAC 660

Qy   661 agtgaattaggcagttgaccaatgttccgcaacacctcccgtctataccaggaaacacaaa 720
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Db   661 AGTGAATTAGGCAGTTGACCCATGTTCCGCAACACCTCCCGTCTATACCAGGGAACACAAA 720
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Qy   721 attgactgggctaagcctggactttcaaggggaaatatgaaaactggagagcaaacaaa 780
         |||||||||||||||||||||||||||||||||||||| |   |  ||||||||||||||
Db   721 ATTGACTGGGCTAAGCCTGGACTTTCAAGGGGAAATATGAAGAGCAGCAAACAAAA 780

Qy   781 gacatggttaaaaggccaaccagaacaattgtcgagccttcaaagcacgagtgccctcagca 840
         ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db   781 GACATGGTTAAAAGGCCAACCAGAACAATTGTCGAGCCTTCAAAGCACCAGTGCCCTCGCA 840

Qy   841 gggacctgaggcattttgcctttaggaaggccagtttcttaaggaaatcttaagaaatc 900
         |||||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||
Db   841 GGGACCCCTGAGGCATTTGCCTTTAGGAAGGCCAGTTTTCTTAAGGAAATCTTAAGAAACTC 900

Qy   901 ttgaaagaatcatgaattttaaccattttaagtataaaacaatatgcagatgcaataatcag 960
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Qy   961 tttagacatgggtcccaaatttttataaagtcaggcatcacaaggatgtcccagctcc 1020
         ||||||||||||||||||||||||||||||||||||| |||||||||| |||||||||||
Db   961 TTTAGACATGGGTCCCAAATTTTTATAAAGTCAGGCATCACAAGGATTGTCCCAGCCTCC 1020

Qy  1021 ggataggtcagaaaacaattagaaaatcaactgtgtccccaatcctaacttttccagaatcag 1080
         |||||||||||||||||| |||||||||||||||||||||||| |||||||||||||||||||
Db  1021 GGATAGGTCAGAAAAACATTAGAAAATCACTGTGTCCCCAATCCTAACTTTTTCCAGAATGATC 1080

Qy  1081 tgtcctagcctcacacaggccccagtcgtctgactctacacacaccacaaccccaa 1140
         ||||||||||||||||||||||| || ||||||||||||||||||||||||||||||
Db  1081 TGTCATAGCCTCCACACAGGCCCGATGTGTCTGACTCTACACACACCACAACCCAA 1140

Qy  1141 gttgcctcaaccattgttaacgtgtcatctcaagtaggtccccattacaaatgccactcccc 1200
         |||||||||||||||| ||||||||||||||||||||||||||||| ||||||||||||||
Db  1141 GTGCCTCAACCATTGTTAACGTGTCATCTCAGTAGGTCCCCATTACAAATGCCACTCCCC 1200

Qy  1201 acctgagctcactgaacctctgctcacaggttcaagcaattctcctgtctcagctcc 1260
         |||||||||||||| |||||||||||||||||||| |||||||||| |||||||||||
Db  1201 ACCTGAGCTCACTGCAACCTCTGCTCACAGGTTCAGCAATTCTCCTGTCTCAGCTCC 1260

Qy  1261 tacagccagaagctccgtgaggtgtctgtcttctacacctaacctgatgtcttac 1320
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Db  1261 TACAGCCAGAAGCTCCGTGAGGGTGTCTGTCTTCACCTACCTGTATGTCTTAC 1320

Qy  1321 acctgagctcactgcaacctctgctccccaggttcaacttctcctgtctcagcctc 1380
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Db  1321 ACCTGAGCTCACTGCAACCTGTGCTCCCCAGTTCAATTCTCCTGTCTCAGCCTCC 1380

Qy  1381 cgcgtaagctgggactacaggcgcgccccggctaatttttgtatttttagtagagatgggg 1440
         ||||||||||||||||||||||||||||||||| ||||||||||||| |||||||||||||
Db  1381 CGCGTAGCTGGGACTACAGGCGCACCACCACGGCTAATTTTGTATTTTTAGTAGAGATGGG 1440

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Db  1441 GTTTCACCATATTAGCCAGGCTGGGTCTTGAACTCCTGACCTCAGGTGATCCACCCTC 1500

Qy  1501 agcctccctaaagtgctgggattaccaggcatgagtccacccgcccaagggtcaggtgt 1560
         |||||||||||||||||||||||||||||||||| ||||||| |||||||||| | ||||
Db  1501 AGCCTCCTAAAGTGCTGGGATTACAGGCGTGAGCCATCACCGCCCAAGGGTCAGTGT 1560

Qy  1561 ttaataagggaatttgaatggtttactaaaacaacagggaaaacaaaagctgtga 1620
         ||||||||||||||| ||||||||||||||||||| ||||||||||||||||||||
Db  1561 TTAATAAGGAATAACTTGAATGGTTTACTAAACAACAGGGAAAACAAAAGCTGTGA 1620

Qy  1621 taattcaaggatggggaatggtgcatggctgctgccctgagtccaggac 1680
         ||||||||||||||||| ||| ||||| || |||||||||||||||| ||
Db  1621 TAATTCAAGGGATTGGGGAATGGTGCCATGAGCTGCCTGCCTAGTCCCGAC 1680

Qy  1681 cactggtcccatcacttcttcccgctaagttcccattcatttatt 1740
         |||| ||||||||||||||||||| |||||| |||||||||||| |
Db  1681 CACTGGTCCTCATCACTTTCTTTCCCTCACTAAAGTCAGCTAAGTTACCATTTTATT 1740

Qy  1741 caccatgctttttgttggtaagcctccacaatcgttactgaaattaagagtatacataactag 1800
         |||||||||| ||||| ||||||||||||||||| |||||||||| |||||||||||||||||
Db  1741 CACCATGCTTTTTGTTGGTAAGCCTCCACAATCGTTACTGAAATTAAGAGTATACATAAACTAG 1800
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Qy   1801 tccattggggccatctgtgtgtgtgtaggggaggggggcatacccagagagactcct 1860
          ||||||||||||||| ||||||||||||||||||| ||| ||||||||||||| ||||
Db   1801 TTCCATTTGGGGCCATCTGTGTGTGTGTATAGGGAGGAGGGCATACCCAGAGAACTCCT 1860

Qy   1861 tgaagcccccggagcagaggtttcctctctgtgtgtgtgtgtcaagagcacccgggtcc 1920
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1861 TGAAGCCCCGGAGCAGAGGTTTCCTCTCTGCGTGTGTGTGTCAAGCTGCACCCCGGGTCC 1920

Qy   1921 tgggtgtgtcctgagcaactcgcagcccgtgccactgttggtttgttatcactctctagg 1980
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1921 TGGGTGTCCTGAGCAACTCGCAGCCGTGCCACTGGGTGTTTGTTTGTTATCACTCTCTAGG 1980

Qy   1981 gaactgttgctttctcatattctgttgactcgttccatccaggcattccatgacaatt 2040
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1981 GACCTGTTGCTTTCTCTATTTCTGTTGACTCGTTCCATCCAGGCATTCATTGACAATT 2040

Qy   2041 tattgagtacttatatctgccagacacacagagaacagagagcaaaaatgttcactgc 2100
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2041 TATTGAGTACTTATATCTGCCAGACACCAGAGAACAGAGAGCAAAAATGTCACTGC 2100

Qy   2101 cctacccttcgtggaggtgacagttccatggaagacgttcagagaaaaattaatagca 2160
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2101 CCTACCTTCGTGGAGGTGACAGTTTCTGTTCCATGGAAGACGTTCATTGACAATT 2160

Qy   2161 gccaacttaaaccagtgctgagaaagaaagaaaagtcactgc 2220
          |||||||||||||||||||||||||||||||||||||||||
Db   2161 GCCAACTTAAAACCCAGTGCTGAAAGAAAAGGAATAAACACCA 2220

Qy   2221 agcatcccttaacaaggccaacctctcctctagcggcccccttgctgtgtccgg 2280
          |||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2221 AGCATCCCTTAACAAGGCCACCTCCTCCTCAGGCGGCCCCTTGCTGTGTGCCCGGG 2280

Qy   2281 ccccaggagtcttccaagccctcctcccacagcgctgcagctggcct 2340
          |||||||||||||||||||||||||||||||||||||||||||||
Db   2281 CCCCCCAAGCCCAGTCTTCCCAAGCCCTCCTCCCATCGTCACAGCCTGCCT 2340

Qy   2341 gcctcgcttcccgtgaatcgtcctggtgcatctgagctgcttggctccaggct 2400
          |||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2341 GCCTCGGCTTCCGTGAATCGTCCATCTGAGCTGGAGCTCCTGGCTCCAGGCT 2400

Qy   2401 ccagaaaggaaatgttgagaggaaactagtctaaacgggaggggacaagttccc 2460
          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2401 CCAGAAAGGAAATGGAGAGGAAACTAGTCTAAGCGGAGGGGGAGACAGTGTTTC 2460

Qy   2461 ctcagaggggaaggggcctccagtccagatcagtcacagcgctgcagcggcct 2520
          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2461 CTCAGAGGGAAAAGGGGCCTCCACGTCCGAGGAAATTCCAGAGGTGCAGCAGGGAG 2520

Qy   2521 tggggaacgctggggctcggggtgctgaagagcaggggtgaaaggggcaggctgaa 2580
          |||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2521 TGGGGACGGCTGGGGCTGCTGAAAGGGCAGGAGAGGGGCAGGGGCAGGCTGAA 2580

Qy   2581 gctgcagagttcagtgtgttcaagtgtggggtgatgcggttc 2640
          |||||||||||||||||||||||||||||||||||||||||
Db   2581 GCTGCCAGAGATGTTCAGTGTGTTCAGCGGGTGCTGGGGACTTCCTGTGAGC 2640

Qy   2641 ccttttatctttctgcttggaggagaagtctattccagaagagttc 2700
          ||||||||||||||||||||||||||||||||||||||||||||||
Db   2641 CTTTTTATCTTTTCTGCTTGGAGGAGAAGTCTATTTCATTGAAGGGAGCAGTTTC 2700

Qy   2701 ataaagtcagctgtcaaaattccagggtgtgcatgggtttccctcaacggaaggcct 2760
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2701 ATAAAGTCAGCTGTTAAAATTCAGGGTGTCCATGGGTTTTCCTTCACGAAGGGCTTTAT 2760

Qy   2761 ttaatggggaatataggaagacagagctccaattccaggatgac 2820
          ||||||||||||||||||||||||||||||||||||||||||||
Db   2761 TTAATGGGGAATATAGGAAGCAGAGCTCCAATTTCCTAGGCCGTTAATTCACGGAGAAGTGAC 2820

Qy   2821 tgggtctttcttcatgtcttctgggggcacttgctggtcggagcaagtctta 2880
          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2821 TGGAGTCTTTTCTTTCATGTCTTCTTCGGGCAACTACTCAGCCCTGGGTGCAACTACTTCAGCCCTTA 2880
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Qy   2881 tgcaagacggtcgaaaaaccttggaatcagaggaggactcggtttctcttctggcttctgccatt 2940
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2881 TGCAAGACGGTCGAAAAACCTTGGAATCAGAGGAGGACTCGGTTTTCTTTCTTTCTGGCCATT 2940

Qy   2941 ggttggctgtgcgaccgtgggcaagtgtctcctcttcctttccctgggggcctagtcttctgct 3000
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2941 GGTTGGCTGTGCGACCGTGGGCAAGTGTCTCCTCTTCCCTGGGGGCCATAGTCTTTCTGCT 3000

Qy   3001 ataaagaaacccttgcagctctcgtgtttctgtgaacacttccctgtgatctctgtagggg 3060
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3001 ATAAAGAAACCCTTGCAGCTCTCGTGTTTCTGTGAACACTTCCCTGTGATCTCTGTAGGGG 3060

Qy   3061 ggatgttgagagtggagggagggcagagactgagccacagagagtgggg 3120
          |||||||||||||||||||||||||||||||||||||||||||||||||
Db   3061 GGATGTTGAGAGTGGAGGGAGGGCAGAGACTGAGCCACAGAGAGTGGGG 3120

Qy   3121 ggacagggaggcaggcagagagctgggtgctccaagcctcagttctccaggacctc 3180
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3121 GGACAGGGAGGCAGGCAGCGAGAGCTGGGTGCTCCAGGCTGGGGTTCTCACTGCTCAGACTC 3180

Qy   3181 caggacccagagagccacaatgctttcaggaaaagctgaaacccaaacagcccacattttcct 3240
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3181 CAGGACCGACAAGCCACAATCCTTCAGGAAAGCTCAATGAACCCAAACGCCACATTTTCCT 3240

Qy   3241 tccctaagcatagaccaagtggcattcaaataacaaaagatgcagagagactggt 3300
          |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3241 TCCCTAAGCATAGACCAAGTGGCATTTCAATAACCAAAAGATGCAGAGAGAGACTGGT 3300

Qy   3301 ggtagcttgctggtcattcaaggaaagctcaagagcagggcaaagtgccagaagattg 3360
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3301 GGTAGCTTTTTGCCTGGCATTCAAAAACTGGGTCGGGCAGAGGCAAAGTGCCAGAGGGG 3360

Qy   3361 ttaaacttttcacctgaccagcgaccacgcccagcgcaggcaagttgacactcacgg 3420
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3361 TTAAACTTTTCACCCTGACCAGCCACCCACGCGCAGGCAAGTTGACTGCTGACACGG 3420

Qy   3421 agtgacctgcagtgcagccagtggagaagaagaagaagagagtagtgtatgagaaag 3480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3421 AGTGACCTGCAGTGCAGCCAGTGGAGAAGAGAAGAAGAGAGTAGTGTATGAGACTTGGT 3480

Qy   3481 acagattcattcaaggggaggctggggaattgaccaacaaaaagagagaaatgcagaaag 3540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3481 ACAGATTCATTCAAGGCGAGGGCAGTGGGAATTGACCACAGGGGATTATATGTCCACGTGATCTGG 3540

Qy   3541 gttctaggaggaggctatatgtggggggaaaaatcagttcaaggaaagtcacaag 3600
          |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3541 GTTCTAGGAGGCAGGGCTATATTGTGGGGGGAAAATCAGTTCAAGGGAAGTCGGGAGA 3600

Qy   3601 cctgaattcctaatacctatattttccctttacaagctgagtaattctgagtcacaag 3660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3601 CCTGATTTCTAATACTATATTTTTCCTTTACAAGCTGAGTAGTTCTGAGTCACAAG 3660

Qy   3661 gtagtaactgaggctgtaagattaactactttaggagtaattctggaacttcctaagag 3720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3661 GTAGTAACTGAGGCTGTAAGATTACTTAGTTGTCAAGGGAGAAACATTCTTTTTCTGT 3720

Qy   3721 gggagttaagcgcacagagggcaatcccgtttctttaacaggaggaagaaacattcctaagag 3780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3721 GGAGTTACCAGCACAAGGGCAATCCCGTTTTCTTTTAACAGGAAGAAAACATTCCTAAGAG 3780

Qy   3781 taaagtccaaacagattcaagcctaggtcttgctgactatagatgagtttttttgaaaat 3840
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3781 TAAAGCCAAACAGATTCAAGCCTAGGTCTTGCTGACTATATGGTTTTTTTTGAAAAAT 3840

Qy   3841 catttcagcgatgtttactactgattactagtaccctttggtcgctg 3900
          |||||||||||||||||||||||||||||||||||||||||||||||
Db   3841 CATTTCAGCGATGTTTACTATCTGATTCAGAAAATGAGACTAGTACCCTTTGGGTCAGCTG 3900

Qy   3901 taaacaaaacaccccagttgtaaatgtctcaaggcttaactgcagaaccaatcaa 3959
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3901 TAAACAAAACACCCCATTTGTAAATGTCTCAAGGTCAAGTTCAGAACCAAATCAAAT 3960

Qy   3960 aagaatagaactctttagagcaaactgtgttctctcccacatctggaggtgagtctgcaggg 4019
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Qy   3961 AAGAATAGAAATCTTTAGAGCAAACTGTGTTTCTCCAC-TCTGGAGGTGAGTCTGCCAGG
          |||||||||||||||||||||||||||||||||||||| |||||||||||||||||||||
Db   4020 CAGTTTGGAAATATTTACTTCACACAAGTATTGACACTGTTGTTCGTATTAACACATAAG 4019

Qy   4020 CAGTTTGGAAATATTTACTTCACACAAGTATTGACACTGTTGTTGGTATTAACACATAAG
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4080 TTGCTCAAAGGCAATCATTATTTCATTATTTCCAAGTGGCTTAAAGTTACTTCTGACAGTTTGGTATA 4079

Qy   4080 TTGCTCAAAGGCAATCATTATTTCATTATTTCCAAGTGGCTTAAAGTTACTTCTGACAGTTTGGTATA
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4140 TTTATTACTATTATTACAGTTGTTGCCAGATACGTTGTTGAAATATTTTATACTTCAAAACT 4139

Qy   4140 tttattggctattgccatttgcttttgtcttttgtttgttttattaatgtaaagca
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Db   4200 TTTATTGGCTATTGCCATTTGCTTTTGTCTTTTGTTTGTTTTATTAATGTAAAGCA 4199

Qy   4200 gggattattaacctacagtccgaaagacctgtgaaattgaatgaggaaaaattacattt
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4260 GGGATTATTAACCTACAGTCCGAAAGACCTGTGAAATTTGAATGAGGAAAAATTACATTT 4259

Qy   4260 tcgtttttaccaccttctctaactaaaatttaactaaacattttattccattgcgaatagagccataa
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4320 TTGTTTTTACCACCTTCTAACTAAAATTTAACTAAACATTTTATTCCATTGCGAATAGAGCCATAA 4319

Qy   4320 acccaaagtggtaataacgtacctgtgatttgtcattaccaatgacacagacat
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4380 ACTCAAAGTGGTAATAACAGTAACCTGTGATTTGTCATTACCAATGACACAGACAT 4379

Qy   4380 ttataactatatatacagtgttgtgcagatacgttgtaagtgaaatatttatatctcaaaact
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db   4500 ACTTTGAAATTAGACCTCCTGCTGAATCTTGTTCTTTTAACATATTAATATAAAAACATGTTTAA 4499

Qy   4500 aatttgatattttgataatcatatatttcattctttgttcctttgtctaatctatatt
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Db   4560 AATTTGATATTTTGATAATCATATTTTCATTATCATTTGTAATCTATATTT 4559

Qy   4560 tatatatattatacacatcttttctgaggaagagttcctccagaattcaccaatgaggttcttg
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Db   4620 TATATATTTGAAACATACTCTTTGTCAGAAGAGTTCCCAGAATTTCACCAATGAGGTTCTTG 4619

Qy   4620 gcatgcaccacacagagtaagaactgatttagaggctaacactttgcctgag
          |||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4680 GCATGCACCACACAGAGTAGGAACTGATTTAGAGGCTAACACTTTGCCTGAG 4679

Qy   4680 atgcaagactgagaagttctctcccaaagatacaagtgtgtttaaagtctaggggt
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Db   4740 ATGCAAGACTGAAATTAGAAAGTTCTCCCAAAGATACAACAGTTGTTTAAAGTCTAGGGGT 4739

Qy   4740 gaggggggaaatctgccgcttctatagaaatgtctctcctggagacctggtaggtgtctgt
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Db   4800 GAGGGGGGAAATCTGCCCGCTTCTATAGGAATGCTCTCCTGGACCCTGGTAGGGTGCTGT 4799

Qy   4800 ccctgtgttctggctggctgtgttattttctctgctctgtccctacagtcttaaaagggttgtt
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   4860 CCTTGTGTTCTGGCTGGCTGTGTTATTTTCTCTGCTCTGTCCCTACGGTCTTAAAGGACTTGTT 4859

Qy   4860 tggatctccagtcctagcaatagtgcctggcacagtgcaggtgttctcaatggagttgcaga
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Db   4920 TGGATCTCCAGTTCCTACGCATAGTGCCTGGCACAGTGCAGGTTCTCAATGAGTTGCAGA 4919

Qy   4920 gtgtaagtgtgtgtacgtgtgtgtgtgtgtgtgtgtgaaatatatatcctgttgaaaaacccaggtggagat
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Db   4980 GTGTAAGTGTGTGTACGTGTGTGTGTGTGTGTGTGAAATATATCCTGTTGAAAAACCCAGGTGGAGAT 4979

Qy   4980 gtgtaagtgtgtacgtgtgtgtgtgtgtgtgtgtgtggtatgggtcataaatttggggtattgggtat
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5040 GTGTAAGTGTGTGTACGTGTGTGTGTGTGTGTGTGTGGATGGTTTCTTTTTAAAAAGAAACT 5039

Qy   5040 ataggaactattattgggtatgggtgcataaatttgaaaact
          |||||||||||||||||||||||||||||||||||||||||
Db   5100 ATAGGAACTATTATTGGGTGTGGGTGCATAAAATTGGGATGTTGTTCTTTTTTAAAAAGAAACT 5099
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Db   5040 ATAGGAACTATTATTGGGTGTGGGTGCATAAAATTGGGATGTTGTTCTTTTTTAAAAAGAAACT 5099

Qy   5100 ccaaacagactcctggaaggttatttctaagagaatcttgctggacgtggcagggcaacc
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5100 CCAAACAGACTTCCTGGAAGGTTATTTCTAAGAGAATCTTGCTGGACGTGGCAGGGCAACCC 5159

Qy   5160 ccctgtgcaagcgcccacccagcgctcacgtggcacctctgtctcccccatgaagggct
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5220 CCCTGTGCACAGCCCCAGCCCTCACGTGGCCACCTCGTCTCTGTCTCCCCATGAAGGGCT 5219

Qy   5220 ggctccccagtatatataaacctctctgggagctcgggcatgagccagcaagggccaccca
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   5280 GGCTCCCCAGTATATATAAAACCTCTCTGGGAGCTCGGGCATGAGCCAGCAAGGGCCACCAT 5279

Qy   5280 ccaggcaacctctcagcacagc
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Db   5280 CCAGGCAACCTCTCAGCACAGC 5300
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RESULT   3
US-08-669A-2
;  Sequence 2, Application US/08938669A
;  Patent No. 6171788
;  GENERAL INFORMATION:
;    APPLICANT: Nguyen, Thai D.
;    APPLICANT: Polansky, Jon R.
;    TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS,
;    TITLE OF INVENTION: PROGNOSIS AND TREATMENT OF GLAUCOMA AND
;    TITLE OF INVENTION: RELATED DISEASES
;    NUMBER OF SEQUENCES: 32
;    CORRESPONDENCE ADDRESS:
;    ADDRESSEE: Howrey & Simon
;    STREET: 1299 Pennsylvania Avenue, N.W.
;    CITY: Washington
;    STATE: DC
;    COUNTRY: USA
;    ZIP: 20004-2402
;    COMPUTER READABLE FORM:
;    MEDIUM TYPE: Diskette
;    COMPUTER: IBM Compatible
;    OPERATING SYSTEM: DOS
;    SOFTWARE: FastSEQ for Windows Version 2.0
;    CURRENT APPLICATION DATA:
;    APPLICATION NUMBER: US/08/938,669A
;    FILING DATE:
;    CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;    APPLICATION NUMBER: 08/791,154
;    FILING DATE: 28-JAN-1997
;    ATTORNEY/AGENT INFORMATION:
;    NAME: Mendelson, Elliot
;    REGISTRATION NUMBER: P-42,878
;    REFERENCE/DOCKET NUMBER: 07425-0034
;    TELECOMMUNICATION INFORMATION:
;    TELEPHONE: 202 383-6857
;    TELEFAX: 202 383-6610
;    TELEX:
;    INFORMATION FOR SEQ ID NO: 2:
;    SEQUENCE CHARACTERISTICS:
;    LENGTH: 5304 base pairs
;    TYPE: nucleic acid
;    STRANDEDNESS: single
;    TOPOLOGY: linear
US-08-938-669A-2
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Query Match             72.6%;  Score 4476;  DB 4;  Length 5304;
Best Local Similarity   99.8%;  Pred. No. 0;
Matches 5016;  Conservative 0;  Mismatches 6;  Indels 2;  Gaps 2;

Qy   1 atcttgttcagtttacctcagggctattatgaaatgagagataaccaatgtgaaag 60
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1 ATCTTTGTTCAGTTTACCTCAGGGCTATTATGAAATGAGATAACCAATGTGAAAG 60
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Qy     61 tcctatataacctgtatagcctccattccggatgtatgtcttcttgggcaggatgaataagaaatcca 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     61 TCCTATATAACCTGTATAGCCTCCATTCCGGATGTATGTCTTCTTGGGCAGGATGAATGAATCA 120

Qy    121 ggagaaggagtatccacgcttagccacgcttagcttgtctgtctcttatttagtga 180
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    121 GGAAGAAGGAGTATCCACGTTAGCCACGCTTGTGTCTGCTCTTATTTAGTGA 180

Qy    181 cagatgttgctcctgacagaagctatctcttcaggaaaactcacatatggtaaatc 240
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    181 CAGATGTTGCTCCTGACAGAAGCTATTCTTCAGGAAAACTCACATATGGTAAATC 240

Qy    241 catcaaaacaggaggaaacaggaatgagatgggcaacttgcccaagagtctcttcc 300
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    241 CATCAAAACAGGAGGAACAGGAATGAGATGGGCAACTTGCCCAAGGAATGCCAG 300

Qy    301 gagagcaaaatagtgatgaaaaatcttcagaaaccttgtttttttaattcagaaaaatg 360
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    301 GAGAGCAAATAATGATGAAAAATCTTCAGAAAACTTTTCCTTTGTTTTTAATTTCAGAAAAATG 360

Qy    361 atgaggaccaaaatcaatgaataaggaaaaacagctcagaaaataggcaag 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    361 ATGAGGACCAAAATCAATGAATAAGGAAAAACAGCTCAGAAAATAGGCAAG 420

Qy    421 ggcatcaactctgggaaggcaagttcagtgacatgtttccaaattgg 480
          |||||||||||||||||||||||||||||||||||||||||||||||
Db    421 GGCATCAACTCTGGGGAGGCAAGTTCAGTGACATGTTTCCAAATTGG 420

Qy    481 agtcattagtattgttcctggggaagacttgcccaaatggggaaa 540
          ||||||||||||||||||||||||||||||||||||||||||||||
Db    481 AGCAAAATCAAAATTCCCAAAATGCAGGGACTGGGAAAGCTTTCATAAC 480

Qy    541 aaacgtcaaaacatgatctgtcagtccccaaaagtcccatccttgtttttaattcaaaaaatg 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    541 AAACGTCAAAACATGATCTGTCAGTCCCCAAAGTCCCATTGTTTTTAATTCAGAAAAATGGGGAA 480

Qy    601 taattaagtaattgttcctgggga 660
          ||||||||||||||||||||||||
Db    601 AGCAAAATCAAAATTCCCAAAATGCAGGGACTGGGAAAGCTTTCATAAC 660

Qy    661 agtggattaagggaggttgacagttgcaaccacctcccgctatacaggggaacacaaaa 720
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    661 AGTGATTAGGCACTTGACCATGTTCGCAACACCTCCCCGTCTATACCAGGGAACACAAA 720

Qy    721 attgactgggctaagcctggacttcaaggggaaatatgaagacaacaaa 780
          |||||||||||||||||||||||||||||||||||||||||||||||||
Db    721 ATTGACTGGGCTAAGCCTGGACTTCAAGGGGAAATATGAAGCAGAGCAAAACAAA 780

Qy    781 gacatggtcaaaggcaaccaggaacattgtgagccttcaagactgggaagctttcatgg 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    781 GACATGGTTAAAGGCAACCAGGAACATTGTGAGCCTTCAAAGCCAGAGTGCCCCTCAGCA 840

Qy    841 ggggacccctgagggccattgcccttttcagaagtccttaaggaaaactc 900
          |||||||||||||||||||||||||||||||||||||||||||||||||||
Db    841 GGGCACCCTGAGGGCCATTTGCCCTTTAGGAAGGCCAGTTTTTCTTAAGGAAAACTTC 900

Qy    901 tctgaaggtcatgaatttcaaccattctttaagtatataaaacaaatatcggatgcaataacag 960
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    901 TTGAAAGATCATGAATTTTAACCATTTTAAGTATAAAACAAATATCGGATGCAATAATCAG 960

Qy    961 tttagacagtgggtcccaattttataaagtcagggcaacatcaaaggagataacgtgtcccagctcc 1020
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    961 TTTAGACATGGGTCCCAATTTTATAAAGTCAGGCATACAAGGAATAACGTGTCCCAGCTCC 1020

Qy   1021 ggataggtcagaaatcactgtgtcccatcctaactttttcagaaatgatc 1080
          ||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1021 GGATAGGTCAGAAATCATTAGAAATACTGTGTCCCATCCTAACTTTTTCAGAAATGATC 1080

Qy   1081 tgtcataagcctcacacacaggccccgatgtgtctgacctacacaccca 1140
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Db   1081 TGTCATAGCCCTCACACACAGGCCCCGATGTGTCTGACCTACAACCACATTCTACACCCCA 1140
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Qy   1141 gtgcctcaaccattgttaacgtgtcatctcagtctaggtccatttcaccttcccc 1200
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Db   1141, GTGCCTCAACCATTGTTAACGTGTCATCTCAGTCTAGGTCCCATTACAAAATGCCACCTCCC 1200

Qy   1201 tgtgcagcccatcccggctccacaggaagtctccccactctagaacttctgcatcacgagt 1260
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Db   1201 TGTGCAGCCCATCCCGGCTCCACAGGAAGTCTCCCCACTCTAGAACTTCTGCATCACGAGT 1260

Qy   1261 tacagcccagaaagctcccgtgaggtgaggtctgtgtctcacacctgtatgtctac 1320
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1261 TACACCCAGAAGCTCCGTGAGGGTGAGGGTCTGTGTCTTACACCCTACCCTCTATGTCTAC 1320

Qy   1321 acctgagctcactgcaacctctgcctcccaaggtcaagcaaatttctctgtctcc 1380
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Db   1321 ACCTGAGCTCACTGCAAACCTCTGCCTCCCAAGGTTCAAGCAAATTTCTCCTGTCTCAGCCTCC 1380

Qy   1381 cgcgtagctgggactacaggcgcacccgggctaatttttgtatttttagtagatggg 1440
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1381 CGCGTAGCTGGGACTACAGGCGCACGCGCCACCACGCCCGGCTAATTTTTGTATTTTTACTAGAGATGGG 1440

Qy   1441 gtttcaccatattagccccgctggtcttgaactcctgacctcggtgatccacccctc 1500
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1441 GTTTCACCATATTAGCCCGGCTGGTCTTGAACTCCTGACCTCAGGTGATCCACCCACCTC 1500

Qy   1501 agcctcccaaagtgctgggattacaggcatgagccaccgccacccggcccaaggttcagtgt 1560
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Db   1501 AGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCGCGCCCGGCCCAAGGTTCAGTGT 1560

Qy   1561 cgcgtagctgggactacaggcgtacgccacgcccggctaatttttgtatttttactagagatgggg 1620
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Db   1561 TTAATAAGGAATAACTTGAATGTGTTACTACAGGGAAAACAACAGAGGTCTGTGA 1620

Qy   1621 taatttcaggggattcctgggaatggggatggtgccatgggctggtctccagac 1680
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1621 TAATTTCACGGGATTTCTTGGGGGCCATTGGGGAAATGGTGCCATGGGTGCCTCCAGAGCACTCCT 1680

Qy   1681 cactggtcctccatacactcttcctcccatttttcaaggtaaagttacccatttatt 1740
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1681 TAATTCAGGGATTCTTGGGGATGGGGAAATGGTGCCATGGGGCTAGGTGTGCTGCCTCAGTGTGT 1740

Qy   1741 caccacgtctttgtgaagccttccacactcgtactgaaataagagtatataaatag 1800
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1741 CACCAGTGCTTTTGTGGGTAAGCCTTCTCATCACTTTCTCCCCATCACTTTTCAGGCTAAGTTACCATTTATT 1800

Qy   1801 ttccattgggggcatctgttgtgtatagggagggggcataccccagagacttcct 1860
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1801 TTCCATTGGGGCCATCTGTTGTGTGTGACTCGTTCATTCATCCAGTCAGGCATTCATTGACCAGG 1860

Qy   1861 tgaagcccccggacgaggtttcctccctccagctgggggagcccctgcaagctcc 1920
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1861 TGAAGCCCCCGGGACGAGGTTTCCTCCCTCCAGCTGGGGGAGCCCCTGCAAGCTCC 1920

Qy   1921 tgggtgtcctgaggcaacctgccagcctgggggagggggaggggcatcctctctctctagg 1980
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1921 TGGGTGTCCTGAGGCAACCTGCCAGCCTGGGGGAGGGGGAGGGGCATCCTCTCTCTTAGG 1980

Qy   1981 gacctgttgcttctgtattctgtgactcgttcattcatccaggcattcattgacaatt 2040
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Db   1981 GACCTGTTGCCTTCTGTTCTGTGACTCGTTCATTCACCTCAGGTGATCCACCCACCTCT 2040

Qy   2041 tattgagttacttatatctgccagaacacagagagaagaaaaatggtgagcagtcactgc 2100
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Db   2041 TATTGAGTTACTTATATCTGCCAGACACAGAGAGAACAAAATGGTGAGCAGTCACTGC 2100

Qy   2101 cctacctccgtgtgggtgacagtttctcatggaagacgaagaaatattaagacca 2160
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Db   2101 CCTACCTTCGTGTGGGTGACAGTTTCTCATGGAAGACGTGCAGAAGAAATTAATAGCCA 2160

Qy   2161 gccaacttaaacccagtgctgaagaagaaataaacaccatcttgaagaattgtgcgc 2220
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Db   2161 GCCAACTTAAACCCAGTGCTGAAGAAGAAAATAAACACCATCTTGAAGAATTGTGCGC 2220

Qy   2221 agcatccccttaacaggccacctcccctagcgcccctgcctccatccgtgccggagg 2280
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Db  2221  AGCATCCCTTAACAAGGCCAACCTCCCTAGGCGGCTCCATCGTGCCCGGGAG  2280
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Qy  2281  ccccccaagagccccgagtccagccctccctccatcacagcgctgcagtgct  2340
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Db  2281  CCCCCAAGGCCCGAGTCTTCCAAGCAGTCACAGTCACACGGCTCCAGCTGGCCT  2340
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Qy  2401  ccagaaaggagaatggaggaggaaactctggaagtgagagaagtgtttc  2460
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Qy  2461  ctcagagaggaaagggcctccacctccacgtccagaggagaggagaggag  2520
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          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  2521  tggggaccgctggggcgtgagccgggtgctgaaagcagatgctgaaag  2580
          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  2521  TGGGGACCGCTGGGGCCTGACCGGGTGCTGAAAGGCAGGAGGTGGCAAGGCTGAA  2580
          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Qy  2581  gctgccagagtgttcaagtgttgttcaggggctgggagtcttctccctgtgagc  2640
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Db  2581  GCTGCCCAGAGTGTTCAGTGTGTTCACGGGGAGTTTCCGGTTGCTTCCTGTGAGC  2640
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Qy  2641  ctttttatctttctttcatgtcttgggggggagaagaatctattatcagaggagtttc  2700
          ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  2641  CTTTTTATCTTTCTTCTTCATGTCTTGGACAGAAGCTCCACGTCCCAGGGAG  2700
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Qy  2881  tgcaagacggtcgaaaacacttggaaatcaggagagaactcggtgtttttctgccatt  2940
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Qy  3121  ggacaggagaggcaggctgggctgtcccatcagtcaactcaactcagagctc  3180
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Qy  3241  tcccctaagacataagaacatggcattgccaaaaaagaatgcaagaactggt  3300
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Db  3301  GGTAGCTTTTGCCTGGCCATTCAAAAAACTGGGCCCAGCTCAAGCTGACTGGGAAAATGCCAGAGAGATTG  3360
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Qy  3361  ttaaacttttcacctgaccagcacacccccgagctccagctgactgttgctactgaacgaggacggcacgg  3420
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Qy  4020  cagtttggaaatatttaacttaacttcacaagttcttcagagaaaatgagactagtaccccttggctg  4079
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Qy  4260  ttgttttcaaagggcaatcaatcataaatttaacattttcattgcaattagaacocataa  4319
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RESULT 4
US-08-938-669A-26
Sequence 26, Application US/0938669A
Patent No. 6171788
GENERAL INFORMATION:
APPLICANT: Nguyen, Thai D.
APPLICANT: Polansky, Jon R.
TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS,
TITLE OF INVENTION: PROGNOSIS AND TREATMENT OF GLAUCOMA AND
TITLE OF INVENTION: RELATED DISEASES
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Howrey & Simon
STREET: 1299 Pennsylvania Avenue, N.W.
CITY: Washington
STATE: DC
COUNTRY: USA
ZIP: 20004-2402
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/938,669A
PRIOR APPLICATION DATA:
FILING DATE:
CLASSIFICATION: 435
APPLICATION NUMBER: 08/791,154
FILING DATE: 28-JAN-1997
ATTORNEY/AGENT INFORMATION:

NAME: Mendelson, Elliot
REGISTRATION NUMBER: P-42,878
REFERENCE/DOCKET NUMBER: 07425-0034
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202 383-6857
TELEFAX: 202 383-6610
TELEX:
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 1548 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
US-08-938-669A-26

Query Match 10.4%; Score 640; DB 4; Length 1548;
Best Local Similarity 100.0%; Pred. No. 6.1e-286;
Matches 640; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 5301 agagctttccagaggaaggaagcctcaccaaagcctctgcaatgaaggtcttctggtcacgttgc 5360
Db 1 AGAGCTTTCCAGAGGAAGCCTCACCAAAGCCTCTGCAATGAAGGTTCTTCTGGTCACGTTGC 60

Qy 5361 tgcagcttctgggcctgagatgccagctgccaagctctgctcagccctggtctggtgtgg 5420
Db 61 TGCAGCTTTGTGGGCCTGAGATGCCAGCTGCCAAGCTCTGCTCTTCTGGTGTGG 120

Qy 5421 gatgtggggccaggacagctcagctcaggaaggccaatgaccagagtggccgatgccag 5480
Db 121 GATGTGGGGCCAGGACAGCTCAGCTCAGGAAGGCCAATGACCAGAGTGGCCGATGCCAG 180

Qy 5481 tataccttcagtgtgggccagtcccaaatgaatccagctgcccagaagcagaagcccatg 5540
Db 181 TATACCTTCAGTGTGGGCCAGTCCCAAATGAATCCAGCTGCCCAGAAGCAGAAGCCCATG 240

Qy 5541 tcagtcatccataacttacagagagagacagcagccagccacccaacgcttagaccttggaaggccacc 5600
Db 241 TCAGTCATCCATAACTTACAGAGAGAGACAGCAGCCAGCCACCCAACGCTTAGACCTTGGAGGCCACC 300

Qy 5601 aaagctcgactcagctccctgagagcctcctccaccaattgaccttggacagggctgcc 5660
Db 301 AAAGCTCGACTCAGCTCCCTGGAGAGCCTCCTCCACCAATTGACCTTGGACAGGCTGCC 360

Qy 5661 aggccccaggagagacccagagggggctgcagaggagctgggcaccctgaggcgggaacgg 5720
Db 361 AGGCCCCAGGAGAGACCCAGAGGGGGGCTGCAGAGGGGAGCTGGGCACCCTGAGGCGGGAGCGG 420

Qy 5721 gaccagctcggaaaaccaagagggagagagtcggagactgcctcacagcaacctcctccgagac 5780
Db 421 GACCAGCTCGGAAAACCCAAACCAGAGTGGAGAACTGCCTCACAGCAACCTCCTCCGAGAC 480

Qy 5781 aagtcagttcctggaggagaagagaagacgactaaggcaagaaaatgagaaatctggcagg 5840
Db 481 AAGTCAGTTCCTGGAGGAGAAGAGAAGACGACTAAGGCAACTAAGGCAAGAAAATGAGAAATCTGGCCAG 540

Qy 5901 cgagacaactgctcggggctgtgccaccaggctccagagaag 5940
Db 541 AGGTTGGAAAAGCAGCAGCCAGCCAGGAGGTAGCAAGGCTGAGGAAAGGGGCGGCCAGTGTCCCCAGAGACC 600

Db 601 CGAGACAACTGCTCGGGGCTGTGCCACCAGGCTCCAGAGAG 640

RESULT 5
US-08-645-900A-2
Sequence 2, Application US/08645900A
Patent No. 5848879
GENERAL INFORMATION:
APPLICANT: NGUYEN, THAI D.
APPLICANT: POLANSKY, JON R.

Qy 4440 acttgaaattagacctcctgtcgttctgttttaacatattaataaaaacatgttaa 4499
Db 4440 ACTTGAAATTAGACCTCTGCTGGATCTTGTTTTTAACATATTAATAAAAACATGTTTAA 4499

Qy 4500 aatttgatatttgatttgtaatcatatttcattatatctcttgtaatctatatt 4559
Db 4500 AATTTGATATTTGATATTTGATAATCATAATTTCATTATATCTCTTTGTAATCTATATTT 4559

Qy 4560 tatatatattgaaaacatcttttctgagagagagagaggaggttctg 4619
Db 4560 TATATATATTGAAAACATCTTTCTGAGAAGAGAGTTCCCAGAGTTCTTG 4619

Qy 4620 gcatgcacacaccaccagagtaagagaactgattttagaggctaacacttggttgtcctgag 4679
Db 4620 GCATGCACACACCACCAGAGTAAGAGAACTGATTTTAGAGGCTAACACTTGGTTGTCCTGAG 4679

Qy 4680 atgcaagactgaaattagaaagttctcccaaagatacccagttgtttaaagctaggggt 4739
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Qy 4740 gaggggggaaatctgcgccttctatataggaatgctcccctgggagctggtagggttgtgctgt 4799
Db 4740 GAGGGGGGAAATCTGCCGCCTTCTATAGGAATGCTCTCCCTGGGAGCTGGTAGGGTGCTGT 4799

Qy 4800 cctggtgttctggctggctgtattttctctcgtctctacagctcttaaaggacttgtt 4859
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Qy 4860 tggatctccagtcctagcatagtgcctggccagtgtcagttctcaatgagttttgcgaga 4919
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Qy 4920 gtgaatggaaatataacctagaaatatatctcctcttgaaatcagcacaccagtagtcctg 4979
Db 4920 GTGAATGGAAATATAAACTAGAAATATATCTTTGTTGAAATCAGCACACCAGTAGTCCTG 4979

Qy 4980 gtgtaagtgtgtgtacgtgtgtgtgtgtgtgtgtgt 5023
Db 4980 GTGTAAGTGTGTGTACGTGTGTGTGTGTGTGTGTGTGT 5023

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; APPLICANT: HUANG, WEIDONG
; TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS OF
; TITLE OF INVENTION: GLAUCOMA
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: HOWREY & SIMON
; STREET: 1299 PENNSYLVANIA AVE., N.W.
; CITY: WASHINGTON
; STATE: D.C.
; COUNTRY: US
; ZIP: 20004
COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/645,900A
; FILING DATE: 14 MAY 1996
; CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
; NAME: MARSH, DAVID
; REGISTRATION NUMBER: 41,408
TELECOMMUNICATION INFORMATION:
; TELEPHONE: (202) 383-7451
; TELEFAX: (202) 383-6610
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
; LENGTH: 1999 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-08-645-900A-2

Query Match      10.4%;   Score 640;   DB 2;   Length 1999;
Best Local Similarity   100.0%;   Pred. No. 6.1e-286;
Matches 640;   Conservative  0;   Mismatches  0;   Indels  0;   Gaps  0;

Qy 5301 agagcttccagaggaagcctcaccaagccctgcaatgagggttcttctgtgcacgttgc 5360
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Db    1 AGAGCTTTCCAGAGGAAGCCCTCACCAAGCCCTGCAATGAGGTTCTTCTGTGCACGTTGC 60

Qy 5361 tgcagcttgggcctgagatgccagctgtccagctgtccctgcctctgtgtgg 5420
        ||||||| ||||||||||||||||||||||||||||||||||||||||||||
Db   61 TGCAGCTTTGGGCCTGAGATGCCAGCTGTCCAGCTGTCCCTGCCTCTGTGTGG 120

Qy 5421 gatgtggggccagggacagctcagctcaggaaggccaatgaccagagtggccgatgccag 5480
        ||||||||||| ||||||||| |||||||||||||||||||||||||||| ||||||||||
Db  121 GATGTGGGGCCCAGGGACAGCACCTCAGGAAGGCCAATGACCAGAGTGGCCCGATGCCAG 180

Qy 5481 tataccttcagtgtggccagtccaatgaatccagctgccagagagcaggccaggccatg 5540
        |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db  181 TATACCTTCAGTGTGGCCAGTCCAATGAATCCAGCTGCCAGAGAGCAGAGCCAGGCCATG 240

Qy 5541 tcagtcatccataacttacagagacagcagcacccaccacccacc 5600
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Db  241 TCAGTCATCCATAACTTACAGAGAGACAGCAGCACACGCCTTAGACTGGAGGCCACC 300

Qy 5601 aaagctcgactcagctccctgggaggcctccctccaccaattgacctt 5660
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Db  301 AAAGCTCGACTCAGCTCCCTGGACAGCCTTGACCACCAGCTGCC 360

Qy 5661 aaggccccaggagaccgagggggctgcagaagctgagtggggaagcgg 5720
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Qy 5721 gaccagctgggaaacccaaaaccagagagagttggagactg 5780
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Db  421 GACCAGCTGGAAACCCAAACCAGAGAGAGTTGGAGACTGCCTACAGGCAACCTCCTCCGAGAC 480

Qy 5781 aagtcagtcctggaggagagaagcgactaaggcagaaa 5840
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RESULT 7
US-08-667-790A-2
; Sequence 2, Application US/08677790A
; Patent No. 5861497
; GENERAL INFORMATION:
;   APPLICANT: NGUYEN, THAI D.
;   APPLICANT: POLANSKY, JON R.
;   APPLICANT: HUANG, WEIDONG
;   TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS OF
;   TITLE OF INVENTION: GLAUCOMA
;   NUMBER OF SEQUENCES: 3
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE: HOWREY & SIMON
;     STREET: 1299 PENNSYLVANIA AVE., N.W.
;     CITY: WASHINGTON
;     STATE: D.C.
;     COUNTRY: US
;     ZIP: 20004
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE: Floppy disk
;     COMPUTER: IBM PC compatible
;     OPERATING SYSTEM: PC-DOS/MS-DOS
;     SOFTWARE: PatentIn Release #1.0, Version #1.25
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER: US/08/667,790A
;     FILING DATE: 21 JUNE 1996
;     CLASSIFICATION:
;   ATTORNEY/AGENT INFORMATION:
;     NAME: MARSH, DAVID
;     REGISTRATION NUMBER: 41,408
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE: (202) 383-7451
;     TELEFAX: (202) 383-6610
; INFORMATION FOR SEQ ID NO: 2:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH: 1999 base pairs
;     TYPE: nucleic acid
;     STRANDEDNESS: single
;     TOPOLOGY: linear
;   MOLECULE TYPE: cDNA
US-08-667-790A-2
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Query Match             100.0%;  Score 640;  DB 2;  Length 1999;
Best Local Similarity   100.0%;  Pred. No. 6.1e-286;
Matches   640;  Conservative   0;  Mismatches   0;  Indels   0;  Gaps   0;

Qy    181 TATACCTTCAGTGTGGCCAGTCCCAATGAATCCAGCTGCCAGGAGCCAGGCCATG 240
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Db    181 TATACCTTCAGTGTGGCCAGTCCCAATGAATCCAGCTGCCAGGAGCCAGGCCATG 240

Qy    241 TCAGTCATCCATAACTTACAGAGAGACAGCAGCAGCCAAGCGCTTAGACCTGGAGGCCACC 300
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Db    241 TCAGTCATCCATAACTTACAGAGAGACAGCAGCAGCCAAGCGCTTAGACCTGGAGGCCACC 300

Qy    301 AAAGCTCGACTCAGCTCCCTGAGAAGCCTCCTCCACCAATTGACCTTGGACCAGGCTGCC 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    301 AAAGCTCGACTCAGCTCCCTGAGAAGCCTCCTCCACCAATTGACCTTGGACCAGGCTGCC 360

Qy    361 AGCCCCAGGAGGACCCAGGAGGGGGCTGCAGAGGAGGAGGCTGGGGAGGGGCGGGGAGCGG 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    361 AGCCCCAGGAGGACCCAGGAGGGGGCTGCAGAGGAGGAGGCTGGGGAGGGGCGGGGAGCGG 420

Qy    421 GACCAGCTGGAAAACCCAAACCAGAGTTGGACTGCCTACAGCAACCTCCTCCGAGAC 480
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Db    421 GACCAGCTGGAAAACCCAAACCAGAGTTGGACTGCCTACAGCAACCTCCTCCGAGAC 480

Qy    481 AAGTCAGTTCTGAGGAAGAGAGAAGAGAACGACTAAGGCAAGAAAAATGAGAAATCTGGCAGG 540
          |||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    481 AAGTCAGTTCTGAGGAAGAGAGAAGAGAACGACTAAGGCAAGAAAAATGAGAAATCTGGCAGG 540

Qy    541 AGGTTGGAAAGCAGCAGCAGCGACTAGGCTAGACAAGGCCCAGTGTCCCCAGACC 600
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Db    541 AGGTTGGAAAGCAGCAGCAGCGACTAGGCTAGACAAGGCCCAGTGTCCCCAGACC 600

Qy    601 CGAGACACTGCTCGGGGCTGTGCCACCCAGGCTCCAGAGAAG 640
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Db    601 CGAGACACTGCTCGGGGCTGTGCCACCCAGGCTCCAGAGAAG 640
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RESULT 8
US-09-220-459-2
; Sequence 2, Application US/09220459
; Patent No. 6150161
; GENERAL INFORMATION:
;   APPLICANT: Nguyen, Thai D
;   APPLICANT: Polansky, Jon R
;   APPLICANT: Huang, Weidong
;   TITLE OF INVENTION: Methods for the Diagnosis of Glaucoma
;   FILE REFERENCE: 07425.0056
;   CURRENT APPLICATION NUMBER: US/09/220,459
;   CURRENT FILING DATE: 1998-12-24
;   EARLIER APPLICATION NUMBER: US/09/220,459
;   EARLIER FILING DATE: 1997-06-25
;   EARLIER APPLICATION NUMBER: 08/882,238
;   EARLIER FILING DATE: 1996-06-25
;   EARLIER APPLICATION NUMBER: 08/649,432
;   EARLIER FILING DATE: 1996-05-17
;   EARLIER APPLICATION NUMBER: 08/546,568
;   EARLIER FILING DATE: 1995-10-20
;   EARLIER APPLICATION NUMBER: 08/336,235
;   EARLIER FILING DATE: 1994-11-03
;   NUMBER OF SEQ ID NOS: 3
;   SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 2
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;  LENGTH: 1999
;  TYPE: DNA
;  ORGANISM: Homo sapiens
US-09-220-459-2

Query Match      10.4%;    Score 640;   DB 3;   Length 1999;
Best Local Similarity  100.0%;   Pred. No. 6.1e-286;
Matches 640;  Conservative  0;  Mismatches  0;  Indels  0;  Gaps  0;

Qy  5301 agagcttccagagggagaagcctcaccaagcctctgcaatgaggttcttcttgtgcacgttgc 5360
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Db     1 agagcttccagagggagaagcctcaccaagcctctgcaatgaggttcttcttgtgcacgttgc 60

Qy  5361 tgcagcttgggcctgagatgccagctgtgctgctctggcctggcctggtgtgg 5420
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Db    61 tgcagcttgggcctgagatgccagctgtgctgctctggcctggcctggtgtgg 120

Qy  5421 gatgtgggggccaggacagctcagctcaggaaggccaatgaccagagtggccgatgccag 5480
         |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   121 gatgtgggggccaggacagctcagctcaggaaggccaatgaccagagtggccgatgccag 180

Qy  5481 tataccttcagtgtggtggccagtcccaatgaatccagctgtgccagagcagagccaggccatg 5540
         ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   181 tataccttcagtgtggtggccagtcccaatgaatccagctgtgccagagcagagccaggccatg 240

Qy  5541 tcagtcatccataacttacagagagcttagacctggaggccacc 5600
         |||||||||||||||||||||||||||||||||||||||||||
Db   241 tcagtcatccataacttacagagagcttagacctggaggccacc 300

Qy  5601 aaagctcgactcagctctcctggagagcctctccaccaattgacctt 5660
         |||||||||||||||||||||||||||||||||||||||||||||
Db   301 aaagctcgactcagctctcctggagagcctctccaccaattgacctt 360

Qy  5661 aggcccccaggagaccagggggctgcagagaggggaagcgg 5720
         |||||||||||||||||||||||||||||||||||||||
Db   361 aggcccccaggagaccagggggctgcagagaggggaagcgg 420

Qy  5721 gaccagctggaaacccaaaaccagacccagagacttgcag 5780
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Db   421 gaccagctggaaacccaaaaccagacccagagacttgcag 480

Qy  5781 aagtcagttctggaggaagaagaagcgactaaggcagg 5840
         ||||||||||||||||||||||||||||||||||||
Db   481 aagtcagttctggaggaagaagaagcgactaaggcagg 540

Qy  5841 aggttggaaagcagcagccagagcagctgccacc 5900
         ||||||||||||||||||||||||||||||||
Db   541 aggttggaaagcagcagccagagcagctgccacc 600

Qy  5901 cgagacactgctcgggctgtgccaccaggctcccagagaag 5940
         |||||||||||||||||||||||||||||||||||||||
Db   601 cgagacactgctcgggctgtgccaccaggctcccagagaag 640
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RESULT   9
US-08-546-568B-2
Sequence 2, Application US/08546568B
Patent No. 6248867
; GENERAL INFORMATION:
; APPLICANT: NGUYEN, THAI D.
; APPLICANT: POLANSKY, JON R.
; APPLICANT: HUANG, WEIDONG
; TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS OF
; TITLE OF INVENTION: GLAUCOMA
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: HOWREY & SIMON
; STREET: 1299 PENNSYLVANIA AVE., N.W.
; CITY: WASHINGTON
; STATE: D.C.
; COUNTRY: US
; ZIP: 20004

RESULT 10
US-08-645-900A-3
; Sequence 3, Application US/08645900A
; Patent No. 5849879
; GENERAL INFORMATION:
; APPLICANT: NGUYEN, THAI D.
; APPLICANT: POLANSKY, JON R.
; APPLICANT: HUANG, WEIDONG
; TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS OF
; TITLE OF INVENTION: GLAUCOMA
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: HOWREY & SIMON
; STREET: 1299 PENNSYLVANIA AVE., N.W.
; CITY: WASHINGTON
; STATE: D.C.
; COUNTRY: US
; ZIP: 20004
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/645,900A
; FILING DATE: 14 MAY 1996
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:
; NAME: MARSH, DAVID
; REGISTRATION NUMBER: 41,408
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (202) 383-6610
; TELEFAX: (202) 383-7451
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1512 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-08-645-900A-3

Query Match   9.8%; Score 604; DB 2; Length 1512;
Best Local Similarity   100.0%; Pred. No. 2.6e-269;
Matches   604; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

Qy   5337 atgaggttcttctgtgcacgttgctgcacgcttgggcctggagatgccagctgtccagctg 5396
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Qy   5397 ctgctctggcctggtcctggtgtggggaccagcagctcagctcaggaaggc 5456
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Db   61 CTGCTCTGGCCTGGTCCTGGTGTGGGGACCAGCAGCTCAGCTCAGGAAGGCC 120

Qy   5457 aatgaccagagtggccaatgccagtataccttcagtgtggccagtccaatgaatccagc 5516
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Db   121 AATGACCAGAGTGGCCGATGCCAGTATACCTTCAGTGTGGCCAGTCCCAATGAATCCAGC 180

Qy   5517 tgcccagagagcagagcccaggcccatgtcagtcatccataacttacagagagacagac 5576
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Db   181 TGCCCAGAGCAGCAGAGCCAGGCCCATGTCAGTCATCCATAACTTACAGAGAGACAGCAC 240

Qy   5577 caacgcttagacctggaggccaccaaagctcagctcagctcctggagagccctcctccac 5636
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Qy   5637 caattgaccttgaccaggctgccaggcccccagagaccagaccatgggggga 5696
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Qy   5697 ctgggcaccctgagcggggagcgggaaccagctggaaaccaacagagagtggggact 5756
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Db   361 CTGGGCACCCTGAGCGGGGAGCGGGAGGGGGAACCGGGGACCAACGGGGAGACT 420

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Db   421 GCCTACAGCAACCTCCCTCCGGAGACAAGTCAGTTCAGTTCTGGAGGGAAGAGGAAGCG 480

Qy   5817 caagaaaatgagaaatctggccaggaggagttcggaaaacagagcaaccaaggaagcgc 5876
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Db   481 CAAGAAAATGAGAAATCTGGCCAGGAGGAGTTCGGAAAACAGAGCAACCAGAGGGCTG 540

Qy   5877 agaagagggggccagtgtccccagacaccgagacactgctgctcgggctgtgccaccaggg 5936
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Db   541 AGAAGGGGGCCAGTGTCCCCAGACACCGAGACACTGCTCGGGCTGTGCCACCAGGG 600

Qy   5937 gaag 5940
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Db   601 GAAG 604

RESULT 11
US-08-882-238A-3
; Sequence 3, Application US/08882238A
; Patent No. 5854415
; GENERAL INFORMATION:
; APPLICANT: NGUYEN, THAI D.
; APPLICANT: POLANSKY, JON R.
; APPLICANT: HUANG, WEIDONG
; TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS OF
; TITLE OF INVENTION: GLAUCOMA
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: HOWREY & SIMON
; STREET: 1299 PENNSYLVANIA AVE., N.W.
; CITY: WASHINGTON
; STATE: D.C.
; COUNTRY: US
; ZIP: 20004
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/882,238A
; FILING DATE: 25 JUNE 1997
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: MARSH, DAVID
; REGISTRATION NUMBER: 41,408
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (202) 383-6610
; TELEFAX: (202) 383-7451
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1512 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-08-882-238A-3

Query Match   9.8%; Score 604; DB 2; Length 1512;
Best Local Similarity   100.0%; Pred. No. 2.6e-269;
Matches   604; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

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Db   1 ATGAGGTTCTTCTGTGCACGTTGCTGCACGCTTGGGCCTGGAGATGCCAGCTGTCCAGCTG 60

Qy   5397 ctgctctggcctggtgtggggatgtggggccaggacagctcaggacagctcaggaaggcc 5456
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Db    61  CTGCTTCTTGCCTGCCTTGGTGTGGGATGTGGGGGCCAGGACAGCTCAGCTCAGGAAGGCC  120
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Qy  5457  aatgaccagagctggccagtataccttcagtgtggccagtccaatgaatccagc  5516
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Qy  5577  caacgcttagacctggaggccacacaaagctcccctggagaggctccctcac  5636
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Db   361  CTGGGCACCCTGAGCGCGGGAGCGGGACCACAGCTGGAAAACCCCAAAACCAGAGGGAGTTGGAGACT  420

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Db     1  ATGAGGTTCTTCCTGTGCACGTTGCTGCACGTTTGGGCCTGAGATGCCAGCTGTCCAGCTG  60

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Qy  5457  aatgaccagagctggccagtataccttcagtgtggccagtccaatgaatccagc  5516
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Db   121  AATGACCAGAGTGGCCGATGCCAGTATACCTTCAGTGTGGCCAGTCCAATGAATCCAGC  180

Qy  5517  tgcccagagcagagccagcaggccatgtcagtcatctcataactacacttacagagagacagcacc  5576
              ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   181  TGCCCAGAGCAGAGCCAGCCAGCCCATGTCAGTCATCTCATAACTTACAGAGACAGCACAC  240

Qy  5577  caacgcttagacctggaggccacacaaagctcccctggagaggctccctcac  5636
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; EARLIER FILING DATE: 1996-05-17
; EARLIER APPLICATION NUMBER: 08/546,568
; EARLIER FILING DATE: 1995-10-20
; EARLIER APPLICATION NUMBER: 08/336,235
; EARLIER FILING DATE: 1994-11-03
; NUMBER OF SEQ ID NOS: 3
; SOFTWARE: PatentIn Ver. 2.0
;
; SEQ ID NO 3
; LENGTH: 1512
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-220-459-3

Query Match
Best Local Similarity   100.0%;   Score 604;   DB 3;   Length 1512;
Matches   604;   Conservative   0;   Mismatches   0;   Indels   0;   Gaps   0;

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Qy   5397 ctgcttctggcctgcctggttgtggggtgtgggggccaggacagctcagctcaggaaggc 5456
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Qy   5457 aatgaccagagtggccgatgccagtataccttcagtgtggggggccagacagctcagc 5516
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Qy   5517 tgcccagagcagagcccaggccatgtcagtcatccataacttacagagagacaagcacc 5576
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Qy   5577 caacgcttagacctggaggaacctccactggaagtcagctccctcctccac 5636
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Db   241 caacgcttagacctggaggaacctccactggaagtcagctccctcctccac 300

Qy   5637 caattgaccttggaccaggctgccaggagaggaggggctgcaaggctg 5696
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Qy   5697 ctgggcaccctgaggcgggaggggagcccaggaggaggaggtagcaaggctg 5756
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RESULT 14
US-08-546-568B-3
Sequence 3, Application US/08546568B
Patent No. 6248867
GENERAL INFORMATION:
; APPLICANT: NGUYEN, THAI D.
; APPLICANT: POLANSKY, JON R.
; APPLICANT: HUANG, WEIDONG
; TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS OF
; TITLE OF INVENTION: GLAUCOMA

; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: HOWREY & SIMON
; STREET: 1299 PENNSYLVANIA AVE., N.W.
; CITY: WASHINGTON
; STATE: D.C.
; COUNTRY: US
; ZIP: 20004
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/546,568B
; FILING DATE: 20 October 1995
; CLASSIFICATION: 536
; ATTORNEY/AGENT INFORMATION:
; NAME: MARSH, DAVID
; REGISTRATION NUMBER: 41,408
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (202) 383-7451
; TELEFAX: (202) 383-6610
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1512 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
US-08-546-568B-3

Query Match
Best Local Similarity   100.0%;   Score 604;   DB 4;   Length 1512;
Matches   604;   Conservative   0;   Mismatches   0;   Indels   0;   Gaps   0;

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Qy   5457 aatgaccagagtggccgatgccagtataccttcagtgtggggggccagacagctcagc 5516
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Qy   5517 tgcccagagcagagcccaggccatgtcagtcatccataacttacagagagacaagcacc 5576
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Qy   5577 caacgcttagacctggaggaacctccactggaagtcagctccctcctccac 5636
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Qy   5697 ctgggcaccctgaggcgggaggggagccccaggaggaggaggtagcaaggctg 5756
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RESULT 15
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; Sequence 1, Application US/08822999
; Patent No. 6271026
; GENERAL INFORMATION:
; APPLICANT: Stone, Edwin M.
; APPLICANT: Sheffield, Val C.
; APPLICANT: Alward, Wallace L.M.
; TITLE OF INVENTION: GLAUCOMA COMPOSITIONS AND THERAPEUTIC
; TITLE OF INVENTION: AND DIAGNOSTIC USES THEREFOR
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FOLEY, HOAG & ELIOT LLP
; STREET: One Post Office Square
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02109-2170
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/822,999
; FILING DATE: 21-MAR-1997
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/791,347
; FILING DATE: 30-JAN-1997
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/748,479
; FILING DATE: 08-NOV-1996
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/234,218
; FILING DATE: 28-APR-1994
; ATTORNEY/AGENT INFORMATION:
; NAME: Arnold, Beth E.
; REGISTRATION NUMBER: 35,430
; REFERENCE/DOCKET NUMBER: UIA-010.27
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-832-1000
; TELEFAX: 617-832-7000
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2166 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-822-999-1
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Query Match      9.7%;  Score 599;  DB 4;  Length 2166;
Best Local Similarity   99.8%;  Pred No. 5.2e-267;
Matches  649;  Conservative  0;  Mismatches  1;  Indels  0;  Gaps  0;
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Job time: 41014 sec
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